

SEQUENCE LISTING

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<120> METHODS AND MATERIALS RELATING TO METALLOCARBOXYPEPTIDASE-LIKE POLYPEPTIDES AND POLYNUCLEOTIDES

<130> HYS-28

<140> NOT YET ASSIGNED

<141> 2000-09-29

<150> US 09/560,875

<151> 2000-04-27

<150> US 09/496,914

<151> 2000-02-03

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<170> PatentIn version 3.0

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<212> DNA

<213> homo sapiens

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gaattcacgc cagagaatgg attgctcctg ctttttgcca atggttcgtc aaagaaattc      180
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Hys-28

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ctt tat ctt ttg ggg atg ctg gtt cct gga ggg ctg gga tat gat aga 162
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tcc tta gcc caa cac aga caa gag att gtg gac aag tca gtg agt cca 210
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 25 30 35

tgg agc ctg gag acg tat tcc tat aac ata tac cac ccc atg gga gag 258
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atc tat gag tgg atg aga gag atc agt gag aag tac aag gaa gtg gtg 306
 Ile Tyr Glu Trp Met Arg Glu Ile Ser Glu Lys Tyr Lys Glu Val Val
 60 65 70

aca cag cat ttc cta gga gtg acc tat gag acc cac ccc ata tat tat 354
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 Leu Lys Ile Ser Gln Pro Ser Gly Asn Pro Lys Lys Ile Ile Trp Met
 90 95 100

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 Asp Cys Gly Ile His Ala Arg Glu Trp Ile Ala Pro Ala Phe Cys Gln
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 Ile Asp Gly Tyr Ile Tyr Thr Trp Thr Thr Asp Arg Leu Trp Arg Lys
 155 160 165

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 170 175 180

Hys-28

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aaa gct gtt gcc agc ttc ata gag agc aag aag gat gat att ttg tgc Lys Ala Val Ala Ser Phe Ile Glu Ser Lys Lys Asp Asp Ile Leu Cys 220 225 230	786
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20 25 30

Hys-28

Val Asp Lys Ser Val Ser Pro Trp Ser Leu Glu Thr Tyr Ser Tyr Asn
35 40 45

Ile Tyr His Pro Met Gly Glu Ile Tyr Glu Trp Met Arg Glu Ile Ser
50 55 60

Glu Lys Tyr Lys Glu Val Val Thr Gln His Phe Leu Gly Val Thr Tyr
65 70 75 80

Glu Thr His Pro Ile Tyr Tyr Leu Lys Ile Ser Gln Pro Ser Gly Asn
85 90 95

Pro Lys Lys Ile Ile Trp Met Asp Cys Gly Ile His Ala Arg Glu Trp
100 105 110

Ile Ala Pro Ala Phe Cys Gln Trp Phe Val Lys Glu Ile Leu Gln Asn
115 120 125

His Lys Asp Asn Ser Arg Ile Arg Lys Leu Leu Arg Asn Leu Asp Phe
130 135 140

Tyr Val Leu Pro Val Leu Asn Ile Asp Gly Tyr Ile Tyr Thr Trp Thr
145 150 155 160

Thr Asp Arg Leu Trp Arg Lys Ser Arg Ser Pro His Asn Asn Gly Thr
165 170 175

Cys Phe Gly Thr Asp Leu Asn Arg Asn Phe Asn Ala Ser Trp Cys Ser
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Ile Gly Ala Ser Arg Asn Cys Gln Asp Gln Thr Phe Cys Gly Thr Gly
195 200 205

Pro Val Ser Glu Pro Glu Thr Lys Ala Val Ala Ser Phe Ile Glu Ser
210 215 220

Lys Lys Asp Asp Ile Leu Cys Phe Leu Thr Met His Ser Tyr Gly Gln
225 230 235 240

Leu Ile Leu Thr Pro Tyr Gly Tyr Thr Lys Asn Lys Ser Ser Asn His
245 250 255

Pro Glu Met Ile Gln Val Gly Gln Lys Ala Ala Asn Ala Leu Lys Ala
260 265 270

Lys Tyr Gly Thr Asn Tyr Arg Val Gly Ser Ser Ala Asp Ile Leu Tyr
275 280 285

Ala Ser Ser Gly Ser Ser Arg Asp Trp Ala Arg Asp Ile Gly Ile Pro
290 295 300

Phe Ser Tyr Thr Phe Glu Leu Arg Asp Ser Gly Thr Tyr Gly Phe Val
Page 4

Hys-28

305 310 315 320

Leu Pro Glu Ala Gln Ile Gln Pro Thr Cys Glu Glu Thr Met Glu Ala
 325 330 335

Val Leu Ser Val Leu Asp Asp Val Tyr Ala Lys His Trp His Ser Asp
 340 345 350

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Ser Cys Met Ser Leu Leu
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Hys-28

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Thr Asp Arg Leu Trp Arg Lys Ser Arg
35 40

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<210> 8
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Thr His Pro Ile Tyr Tyr Leu Lys Ile
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Hys-28

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<400> 13

Leu Thr Met His Ser Tyr Gly Gln Leu Ile Leu Thr Pro Tyr
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<210> 14
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Gly Thr Asp Leu Asn Arg Asn Phe Asn
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Ile Gly

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Hys-28

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 35 40 45
 Glu Val Val Thr Gln His Phe Leu Gly Val Thr Tyr Glu Thr His Pro
 50 55 60
 Ile Tyr Tyr Leu Lys Ile Ser Gln Pro Ser Gly Asn Pro Lys Lys Ile
 65 70 75 80
 Ile Trp Met Asp Cys Gly Ile His Ala Arg Glu Trp Ile Ala Pro Ala
 85 90 95
 Phe Cys Gln Trp Phe Val Lys Glu Ile Leu Gln Asn His Lys Asp Asn
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 115 120 125
 Val Leu Asn Ile Asp Gly Tyr Ile Tyr Thr Trp Thr Thr Asp Arg Leu
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 Pro Glu Thr Lys Ala Val Ala Ser Phe Ile Glu Ser Lys Lys Asp Asp
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 Ile Leu Cys Phe Leu Thr Met His Ser Tyr Gly Gln Leu Ile Leu Thr
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 Pro Tyr Gly Tyr Thr Lys Asn Lys Ser Ser Asn His Pro Glu Met Ile
 225 230 235 240
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 260 265 270
 Ser Ser Arg Asp Trp Ala Arg Asp Ile Gly Ile Pro Phe Ser Tyr Thr
 275 280 285
 Phe Glu Leu Arg Asp Ser Gly Thr Tyr Gly Phe Val Leu Pro Glu Ala
 290 295 300
 Gln Ile Gln Pro Thr Cys Glu Glu Thr Met Glu Ala Val Leu Ser Val
 305 310 315 320
 Leu Asp Asp Val Tyr Ala Lys His Trp His Ser Asp Ser Ala Gly Arg
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 340 345 350

Hys-28

Leu Leu

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 35 40 45
 Pro Ala Phe Cys Gln Trp Phe Val Lys Glu Ile Leu Gln Asn His Lys
 50 55 60
 Asp Asn Ser Arg Ile Arg Lys Leu Leu Met Asn Leu Asp Phe Tyr Val
 65 70 75 80
 Leu Pro Val Leu Asn Ile Asp Gly Tyr Ile Tyr Thr Trp Thr Thr Asp
 85 90 95
 Arg Leu Trp Arg Lys Ser Arg Ser Pro His Asn Asn Gly Thr Cys Phe
 100 105 110
 Gly Thr Asp Leu Asn Arg Asn Phe Asn Ala Ser Trp Cys Ser Ile Gly
 115 120 125
 Ala Ser Arg Asn Cys Gln Asp Gln Thr Phe Cys Gly Thr Gly Pro Val
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 Tyr Asn Tyr Glu Lys Tyr Asn Ser Trp Glu Lys Ile Asp Ala Trp Thr
 35 40 45
 Ala Asp Ile Ala Asn Glu Asn Pro Ser Leu Val Ser Arg Leu Gln Ile
 50 55 60
 Gly Thr Thr Phe Glu Gly Arg Pro Met Pro Leu Leu Lys Val Gly Lys
 65 70 75 80
 Pro Gly Val Asn Lys Lys Ala Ile Phe Ile Asp Cys Gly Phe His Ala
 85 90 95
 Arg Glu Trp Ile Ser Pro Ala Phe Cys Gln Trp Phe Val Arg Glu Ala
 Page 9

Hys-28

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 Val Arg Thr Tyr Gly Lys Glu Thr Ile Met Thr Gln Leu Leu Asn Lys
 115 120 125
 Leu Asp Phe Tyr Ile Leu Pro Val Leu Asn Ile Asp Gly Tyr Val Tyr
 130 135 140
 Ser Trp Lys Gln Ser Arg Met Trp Arg Lys Thr Arg Ser Val Asn Ala
 145 150 155 160
 Gly Ser Thr Cys Ile Gly Thr Asp Pro Asn Arg Asn Phe Asp Ala Ala
 165 170 175
 Trp Cys Ser Val Gly Ala Ser Arg Asn Pro Cys Ser Glu Thr Tyr Cys
 180 185 190
 Gly Ser Lys Pro Glu Ser Glu Lys Glu Thr Lys Ala Leu Ala Asp Phe
 195 200 205
 Ile Arg Arg Asn Arg Ser Ile Ile Gln Ala Tyr Leu Thr Ile His Ser
 210 215 220
 Tyr Ser Gln Met Leu Leu Tyr Pro Tyr Ser Tyr Thr Tyr Asp Leu Thr
 225 230 235 240
 Ser Asn Asn Lys Lys Leu Asn Ser Ile Ala Lys Glu Ala Ile Arg Glu
 245 250 255
 Leu Lys Val Leu Phe Gly Thr Glu Tyr Thr Tyr Gly Pro Gly Ala Ala
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 Thr Ile Tyr Pro Ala Ala Gly Gly Ser Asp Asp Trp Ala Tyr Asp Gln
 275 280 285
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 20 25 30
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 35 40 45
 Ala Gly Gln Asn Lys Pro Ala Ile Phe Met Asp Cys Gly Phe His Ala
 50 55 60
 Arg Glu Trp Ile Ser Pro Ala Phe Cys Gln Trp Phe Val Arg Glu Ala
 65 70 75 80
 Val Arg Thr Tyr Gly Arg Glu Ile Gln Val Thr Glu Leu Leu Asp Lys
 85 90 95
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 Page 10

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